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RAW SEQUENCE LISTING   DATE: 06/05/2001
PATENT APPLICATION: **US/09/804,733A**   TIME: 13:00:38

Input Set : **A:\Mtc661~1.txt**
Output Set: **C:\CRF3\06052001\I804733A.raw**

ENTERED

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 3 <110> APPLICANT: Monsanto Company
 5 <120> TITLE OF INVENTION: RECOMBINANT PROTEINS CONTAINING REPEATING UNITS
 7 <130> FILE REFERENCE: MTC6614.2
 9 <140> CURRENT APPLICATION NUMBER: US 09/804,733A
10 <141> CURRENT FILING DATE: 2001-03-13
12 <150> PRIOR APPLICATION NUMBER: US 60/188,990
13 <151> PRIOR FILING DATE: 2000-03-13
15 <160> NUMBER OF SEQ ID NOS: 29
17 <170> SOFTWARE: PatentIn version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 5
21 <212> TYPE: PRT
22 <213> ORGANISM: Euthynnus pelamis
24 <220> FEATURE:
25 <221> NAME/KEY: PEPTIDE
26 <222> LOCATION: (1)..(5)
28 <400> SEQUENCE: 1
30 Leu Lys Pro Asn Met
31 1               5
33 <210> SEQ ID NO: 2
34 <211> LENGTH: 4
35 <212> TYPE: PRT
36 <213> ORGANISM: Euthynnus pelamis
38 <220> FEATURE:
39 <221> NAME/KEY: PEPTIDE
40 <222> LOCATION: (1)..(4)
42 <400> SEQUENCE: 2
44 Lys Pro Asn Met
45 1
47 <210> SEQ ID NO: 3
48 <211> LENGTH: 4
49 <212> TYPE: PRT
50 <213> ORGANISM: Euthynnus pelamis
52 <220> FEATURE:
53 <221> NAME/KEY: PEPTIDE
54 <222> LOCATION: (1)..(4)
56 <400> SEQUENCE: 3
58 Val Val Tyr Pro
59 1
61 <210> SEQ ID NO: 4
62 <211> LENGTH: 15
63 <212> TYPE: DNA
C--> 64 <213> ORGANISM: Artificial
66 <220> FEATURE:
67 <221> NAME/KEY: misc_feature
68 <222> LOCATION: (1)..(15)
69 <223> OTHER INFORMATION: Degenerate sequence
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       73 <221> NAME/KEY: misc_feature
       74 <222> LOCATION: (1)..(15)
       75 <223> OTHER INFORMATION: n=a, t, c or g; r=a or g; y=c or t
       78 <400> SEQUENCE: 4                                                        15
W-->   79 ctnaarccna ayatg
       82 <210> SEQ ID NO: 5
       83 <211> LENGTH: 60
       84 <212> TYPE: DNA
C-->   85 <213> ORGANISM: Artificial
       87 <220> FEATURE:
       88 <221> NAME/KEY: misc_feature
       89 <222> LOCATION: (1)..(60)
       90 <223> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t
       93 <220> FEATURE:
       94 <221> NAME/KEY: misc_feature
       95 <222> LOCATION: (1)..(60)
       96 <223> OTHER INFORMATION: Degenerate sequence
       99 <400> SEQUENCE: 5
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      103 <210> SEQ ID NO: 6
      104 <211> LENGTH: 60
      105 <212> TYPE: DNA
C-->  106 <213> ORGANISM: Artificial
      108 <220> FEATURE:
      109 <221> NAME/KEY: misc_feature
      110 <222> LOCATION: (1)..(60)
      111 <223> OTHER INFORMATION: n=any nucleotide, r=a or g, y=c or t
      114 <220> FEATURE:
      115 <221> NAME/KEY: misc_feature
      116 <222> LOCATION: (1)..(60)
      117 <223> OTHER INFORMATION: degenerate sequence
      120 <400> SEQUENCE: 6
W-->  121 catrttnggy ttnagcatrt tnggyttnag catrttnggy ttnagcatrt tnggyttnag      60
      124 <210> SEQ ID NO: 7
      125 <211> LENGTH: 25
      126 <212> TYPE: DNA
C-->  127 <213> ORGANISM: Artificial
      129 <220> FEATURE:
      130 <221> NAME/KEY: misc_feature
      131 <222> LOCATION: (1)..(25)
      132 <223> OTHER INFORMATION: Primer
      135 <220> FEATURE:
      136 <221> NAME/KEY: misc_feature
      137 <222> LOCATION: (1)..(25)
      138 <223> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t
      141 <400> SEQUENCE: 7                                                       25
W-->  142 aaagaattcc tnaarccnaa yatgc
      145 <210> SEQ ID NO: 8
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RAW SEQUENCE LISTING
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Input Set : A:\Mtc661~1.txt
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    152 <222> LOCATION: (1)..(27)
    153 <223> OTHER INFORMATION: Primer
    156 <220> FEATURE:
    157 <221> NAME/KEY: misc_feature
    158 <222> LOCATION: (1)..(27)
    159 <223> OTHER INFORMATION: n=any nucleotide; r=a or g; y=c or t
    162 <400> SEQUENCE: 8
W--> 163 aaagcggccg ccatrttngg yttnagc                                    27
    166 <210> SEQ ID NO: 9
    167 <211> LENGTH: 20
    168 <212> TYPE: DNA
C--> 169 <213> ORGANISM: Artificial
    171 <220> FEATURE:
    172 <221> NAME/KEY: misc_feature
    173 <222> LOCATION: (1)..(20)
    174 <223> OTHER INFORMATION: Primer
    177 <400> SEQUENCE: 9
    178 taatacgact cactataggg                                            20
    181 <210> SEQ ID NO: 10
    182 <211> LENGTH: 19
    183 <212> TYPE: DNA
C--> 184 <213> ORGANISM: Artificial
    186 <220> FEATURE:
    187 <221> NAME/KEY: misc_feature
    188 <222> LOCATION: (1)..(19)
    189 <223> OTHER INFORMATION: Primer
    192 <400> SEQUENCE: 10
    193 cgatcaataa cgagtcgcc                                             19
    196 <210> SEQ ID NO: 11
    197 <211> LENGTH: 48
    198 <212> TYPE: DNA
C--> 199 <213> ORGANISM: Artificial
    201 <220> FEATURE:
    202 <221> NAME/KEY: misc_feature
    203 <222> LOCATION: (1)..(48)
    204 <223> OTHER INFORMATION: n=any nucleotide; y=c or t
    207 <220> FEATURE:
    208 <221> NAME/KEY: misc_feature
    209 <222> LOCATION: (1)..(48)
    210 <223> OTHER INFORMATION: Degenerate sequence
    213 <400> SEQUENCE: 11
W--> 214 gtngtntayc cngtngtnta yccngtngtn tayccngtng tntayccn            48
    217 <210> SEQ ID NO: 12
    218 <211> LENGTH: 48
```

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C--> 220 <213> ORGANISM: Artificial
      222 <220> FEATURE:
      223 <221> NAME/KEY: misc_feature
      224 <222> LOCATION: (1)..(48)
      225 <223> OTHER INFORMATION: n=any nucleotide; r=a or g
      228 <220> FEATURE:
      229 <221> NAME/KEY: misc_feature
      230 <222> LOCATION: (1)..(48)
      231 <223> OTHER INFORMATION: Degenerate sequence
      234 <400> SEQUENCE: 12
W--> 235 nggrtanacn acnggrtana cnacnggrta nacnacnggr tanacnac                48
      238 <210> SEQ ID NO: 13
      239 <211> LENGTH: 33
      240 <212> TYPE: DNA
C--> 241 <213> ORGANISM: Artificial
      243 <220> FEATURE:
      244 <221> NAME/KEY: misc_feature
      245 <222> LOCATION: (1)..(33)
      246 <223> OTHER INFORMATION: Forward primer
      249 <220> FEATURE:
      250 <221> NAME/KEY: misc_feature
      251 <222> LOCATION: (1)..(33)
      252 <223> OTHER INFORMATION: n=any nucleotide; y=c or t
      255 <400> SEQUENCE: 13
W--> 256 aaaggatccg tngtntaycc ngtngtntay ccn                            33
      259 <210> SEQ ID NO: 14
      260 <211> LENGTH: 33
      261 <212> TYPE: DNA
C--> 262 <213> ORGANISM: Artificial
      264 <220> FEATURE:
      265 <221> NAME/KEY: misc_feature
      266 <222> LOCATION: (1)..(33)
      267 <223> OTHER INFORMATION: Reverse primer
      270 <220> FEATURE:
      271 <221> NAME/KEY: misc_feature
      272 <222> LOCATION: (1)..(33)
      273 <223> OTHER INFORMATION: n=any nucleotide; r=a or g
      276 <400> SEQUENCE: 14
W--> 277 cccaagcttn ggrtanacna cnggrtanac nac                            33
      280 <210> SEQ ID NO: 15
      281 <211> LENGTH: 45
      282 <212> TYPE: DNA
C--> 283 <213> ORGANISM: Artificial
      285 <220> FEATURE:
      286 <221> NAME/KEY: misc_feature
      287 <222> LOCATION: (1)..(45)
      288 <223> OTHER INFORMATION: n=any nucleotide
      291 <220> FEATURE:
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      292 <221> NAME/KEY: misc_feature
      293 <222> LOCATION: (1)..(45)
      294 <223> OTHER INFORMATION: Degenerate sequence
      297 <400> SEQUENCE: 15                                                45
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      301 <210> SEQ ID NO: 16
      302 <211> LENGTH: 45
      303 <212> TYPE: DNA
C--> 304 <213> ORGANISM: Artificial
      306 <220> FEATURE:
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      308 <222> LOCATION: (1)..(45)
      309 <223> OTHER INFORMATION: n=any nucleotide
      312 <220> FEATURE:
      313 <221> NAME/KEY: misc_feature
      314 <222> LOCATION: (1)..(45)
      315 <223> OTHER INFORMATION: Degenerate sequence
      318 <400> SEQUENCE: 16                                                45
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      322 <210> SEQ ID NO: 17
      323 <211> LENGTH: 36
      324 <212> TYPE: DNA
C--> 325 <213> ORGANISM: Artificial
      327 <220> FEATURE:
      328 <221> NAME/KEY: misc_feature
      329 <222> LOCATION: (1)..(36)
      330 <223> OTHER INFORMATION: Forward primer
      333 <220> FEATURE:
      334 <221> NAME/KEY: misc_feature
      335 <222> LOCATION: (1)..(36)
      336 <223> OTHER INFORMATION: n=any nucleotide
      339 <400> SEQUENCE: 17                                                36
W--> 340 aaaggatccg tnccnccngt nccnccngtn ccnccn
      343 <210> SEQ ID NO: 18
      344 <211> LENGTH: 36
      345 <212> TYPE: DNA
C--> 346 <213> ORGANISM: Artificial
      348 <220> FEATURE:
      349 <221> NAME/KEY: misc_feature
      350 <222> LOCATION: (1)..(36)
      351 <223> OTHER INFORMATION: Reverse primer
      354 <220> FEATURE:
      355 <221> NAME/KEY: misc_feature
      356 <222> LOCATION: (1)..(36)
      357 <223> OTHER INFORMATION: n=any nucleotide
      360 <400> SEQUENCE: 18                                                36
W--> 361 aataagcttn ggnggnacng gnggnacngg nggnac
      364 <210> SEQ ID NO: 19
      365 <211> LENGTH: 8
```

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220> to
<223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/804,733A**

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Input Set : **A:\Mtc661~1.txt**
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L:64  M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:79  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:85  M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:106 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:127 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:142 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:148 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:169 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:184 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:199 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:214 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:220 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:241 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:262 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:283 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:298 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:304 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:325 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:340 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:346 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18
L:361 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:367 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19
L:383 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:20
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:404 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:21
L:419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:425 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:22
L:440 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:446 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:23
L:461 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:469 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
L:485 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
L:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:508 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
L:523 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:525 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:531 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27
L:546 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
```

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```
L:554 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:28
L:569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:571 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:577 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
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